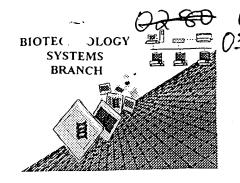
RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09/778, 47/
Source:	OIPE
Date Processed by STIC:	3-1-01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,

2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 c-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 c-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable-form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

OIPE

RAW SEQUENCE LISTING

DATE: 10/30/2001

PATENT APPLICATION: US/09/778,971

TIME: 13:01:02

Input Set : A:\6138SEQ.txt

Output Set: N:\CRF3\10302001\I778971.raw

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1 <110> APPLICANT: Shaughnessy, John D.
 3 <120> TITLE OF INVENTION: Evi27 Gene Sequence and Protein Encoded Thereby
 5 <130> FILE REFERENCE: D6138
 7 <140> CURRENT APPLICATION NUMBER: US 09/778,971
 8 <141> CURRENT FILING DATE: 2001-02-02
10 <150> PRIOR APPLICATION NUMBER: US 60/180, 374
11 <151> PRIOR FILING DATE: 2000-02-04
13 <160> NUMBER OF SEQ ID NOS: 9
16 <210> SEQ ID NO: 1
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17 <211> LENGTH: 1827
18 <212> TYPE: DNA
19 <213> ORGANISM: Homo sapiens
21 <220> FEATURE:
23 <221> NAME/KEY: prim_transcript
24 <223> OTHER INFORMATION: cDNA of human Evi27
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29 gtaccccgag agccgaccgt tcaatgtggc tctgaaactg ggccatctcc
                                                           100
30 agagtggatg ctacaacatg atctaatccc cggagacttg agggacctcc
31 gagtagaacc tgttacaact agtgttgcaa caggggacta ttcaattttg
32 atgaatgtaa gctgggtact ccgggcagat gccagcatcc gcttgttgaa
33 ggccaccaag atttgtgtga cgggcaaaag caacttccag tcctacagct
34 gtgtgaggtg caattacaca gaggccttcc agactcagac cagaccctct
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35 ggtggtaaat ggacattttc ctatatcggc ttccctgtag agctgaacac
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36 agtotatttc attggggccc ataatattcc taatgcaaat atgaatgaag
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37 atggcccttc catgtctgtg aatttcacct caccaggctg cctagaccac
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38 ataatgaaat ataaaaaaa gtgtgtcaag gccggaagcc tgtgggatcc
39 gaacatcact gcttgtaaga agaatgagga gacagtagaa gtgaacttca
                                                           600
40 caaccactcc cctgggaaac agatacatgg ctcttatcca acacagcact
                                                           650
41 atcatcgggt tttctcaggt gtttgagcca caccagaaga aacaaacgcg
                                                           700
42 agcttcagtg gtgattccag tgactgggga tagtgaaggt gctacggtgc
                                                           750
43 agetgaetee atatttteet aettgtggea gegaetgeat eegaeataaa
44 ggaacagttg tgctctgccc acaaacaggc gtccctttcc ctctggataa
45 caacaaaagc aagccgggag gctggctgcc tctcctcctg ctgtctctgc
46 tggtggccac atgggtgctg gtggcaggga tctatctaat gtggaggcac
47 gaaaggatca agaagacttc cttttctacc accacactac tgccccccat 1000
48 taaggttett gtggtttace catetgaaat atgttteeat cacacaattt 1050
49 gttacttcac tgaatttctt caaaaccatt gcagaagtga ggtcatcctt 1100
50 gaaaagtggc agaaaaagaa aatagcagag atgggtccag tgcagtggct 1150
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51 tgccactcaa aagaaggcag cagacaaagt cgtcttcctt ctttccaatg 1200 52 acgtcaacag tgtgtgcgat ggtacctgtg gcaagagcga gggcagtccc 1250 53 agtgagaact ctcaagacct cttccccctt gcctttaacc ttttctgcag 1300 54 tgatctaaga agccagattc atctgcacaa atacgtggtg gtctacttta 1350 55 gagagattga tacaaaagac gattacaatg ctctcagtgt ctgccccaag 1400 56 taccacttca tgaaggatgc cactgctttc tgtgcagaac ttctccatgt 1450 57 caagcagcag gtgtcagcag gaaaaagatc acaagcctgc cacgatggct 1500

58 getgeteett gtageceace catgagaage aagagaeett aaaggettee 1550 filed with 10/7/01 dishertes but not processed until 10/30/0) RAW SEQUENCE LISTING DATE: 10/30/2001 PATENT APPLICATION: US/09/778,971 TIME: 13:01:02

Input Set : A:\6138SEQ.txt

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     61 tttcaaatat tactaactaa tgtagcatta actaacgatt ggaaactaca 1700
     62 tttacaactt caaagctgtt ttatacatag aaatcaatta cagctttaat 1750
     63 tgaaaactgt aaccattttg ataatgcaac aataaagcat cttccaaaaa 1800
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     68 <212> TYPE: DNA
     69 <213> ORGANISM: Homo sapiens
     71 <220> FEATURE:
     73 <221> NAME/KEY: prim_transcript
     74 <223> OTHER INFORMATION: cDNA of human Evi27
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                                                                100
     78 gtaccccgag agccgaccgt tcaatgtggc tctgaaactg ggccatctcc
     79 agagtggatg ctacaacatg atctaatccc cggagacttg agggacctcc
                                                                150
     80 gagtagaacc tgttacaact agtgttgcaa caggggacta ttcaattttg
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     81 atgaatgtaa gctgggtact ccgggcagat gccagcatcc gcttgttgaa
                                                                250
     82 ggccaccaag atttgtgtga cgggcaaaag caacttccag tcctacagct
                                                                300
     83 gtgtgaggtg caattacaca gaggccttcc agactcagac cagaccctct
     84 ggtggtaaat ggacattttc ctatatcggc ttccctgtag agctgaacac
                                                                400
     85 agtotattic attggggccc ataatattcc taatgcaaat atgaatgaag
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     86 atggcccttc catgtctgtg aatttcacct caccaggctg cctagaccac
                                                                500
     87 ataatgaaat ataaaaaaaa gtgtgtcaag gccggaagcc tgtgggatcc
                                                                550
                                                                600
     88 qaacatcact qcttgtaaga agaatgagga gacagtagaa gtgaacttca
     89 caaccactcc cctgggaaac agatacatgg ctcttatcca acacagcact
                                                                650
     90 atcatcgggt tttctcaggt gtttgagcca caccagaaga aacaaacgcg
                                                                700
     91 agcttcagtg gtgattccag tgactgggga tagtgaaggt gctacggtgc
                                                                750
     92 aggtaaagtt cagtgagctg ctctggggag ggaagggaca tagaagactg
                                                                800
     93 ttccatcatt cattgctttt aaggatgagt tctctcttgt caaatgcact
                                                                850
     94 tetgecagea gacaceagtt aagtggegtt catgggggtt etttegetge
     95 agectecace gtgetgaggt caggaggeeg aegtggeagt tgtggteect
     96 tttgcttgta ttaatggctg ctgaccttcc aaagcacttt ttattttcat 1000
     97 tttctgtcac agacactcag ggatagcagt accattttac ttccgcaagc 1050
     98 ctttaactgc aagatgaagc tgcaaagggt ttgaaatggg aaggtttgag 1100
     99 ttccaggcag cgtatgaact ctggagaggg gctgccagtc ctctctgggc 1150
     100 cgcagcggac ccagctggaa cacaggaagt tggagcagta ggtgctcctt 1200
     101 cacctctcag tatgtctctt tcaactctag tttttgaagt ggggacacag 1250
     102 gaagtecagt ggggacacag ceaetececa aagaataagg aacttecatg 1300
W--> 103 cttcattccc tggcataaaa agtgntcaaa cacaccagag ggggcaggca 1350
     104 ccaqccaggg tatgatgggt actaccettt tetggagaac catagaette 1400
     105 ccttactaca gggacttqca tqtcctaaag cactggctga aggaagccaa 1450
     106 gaggateact getgeteett ttttgtagag gaaatgtttg tgtaegtggt 1500
     107 aagatatgac ctagcccttt taggtaagcg aactggtatg ttagtaacgt 1550
     108 gtacaaagtt taggttcaga ccccgggagt cttgggcatg tgggtctcgg 1600
     109 qtcactqqtt ttqactttag ggctttgtta cagatgtgtg accaagggga 1650
     110 aaatgtgcat gacaacacta gaggtagggg cgaagccaga aagaagggaa 1700
     111 gttttggctg aagtaggagt cttggtgaga ttttgctgtg atgcatggtg 1750
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RAW SEQUENCE LISTING DATE: 10/30/2001 PATENT APPLICATION: US/09/778,971 TIME: 13:01:02

Input Set : A:\6138SEQ.txt

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113 cttgtggcag cgactgcatc cgacataaag gaacagttgt gctctgccca 1850
114 caaacaggcg tccctttccc tctggataac aacaaaagca agccgggagg 1900
115 ctggctgcct ctcctcctgc tgtctctgct ggtggccaca tgggtgctgg 1950
116 tggcagggat ctatctaatg tggaggcacg aaaggatcaa gaagacttcc 2000
117 ttttctacca ccacactact gccccccatt aaggttcttg tggtttaccc 2050
118 atctgaaata tgtttccatc acacaatttg ttacttcact gaatttcttc 2100
119 aaaaccattg cagaagtgag gtcatccttg aaaagtggca gaaaaagaaa 2150
120 atagcagaga tgggtccagt gcagtggctt gccactcaaa agaaggcagc 2200
121 agacaaagtc gtcttccttc tttccaatga cgtcaacagt gtgtgcgatg 2250
122 gtacctgtgg caagagcgag ggcagtccca gtgagaactc tcaagacctc 2300
123 ttcccccttg cctttaacct tttctgcagt gatctaagaa gccagattca 2350
124 tetgeacaaa tacgtggtgg tetaetttag agagattgat acaaaagaeg 2400
125 attacaatgc tctcagtgtc tgccccaagt accacttcat gaaggatgcc 2450
126 actgctttct gtgcagaact tctccatgtc aagcagcagg tgtcagcagg 2500
127 aaaaagatca caagcctgcc acgatggctg ctgctccttg tagcccaccc 2550
128 atgagaagca agagacetta aaggetteet ateecaceaa ttacagggaa 2600
129 aaaacgtgtg atgatcctga agcttactat gcagcctaca aacagcctta 2650
131 qtaqcattaa ctaacqattq qaaactacat ttacaacttc aaagctgttt 2750
132 tatacataga aatcaattac agctttaatt gaaaactgta accattttga 2800
2856
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136 <210> SEQ ID NO: 3
137 <211> LENGTH: 1963
138 <212> TYPE: DNA
139 <213> ORGANISM: Unknown
141 <220> FEATURE:
143 <221> NAME/KEY: prim_transcript
144 <223> OTHER INFORMATION: cDNA of mouse Evi27
146 <400> SEOUENCE: 3
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148 gtgcaggagc gccctgcctc gagagccgac tattcagtgt ggctctgaga
                                                          100
149 cagggecate tecaggagtgg atggtecaae acacacteae tecaggagae
150 ttgagggacc tccaagtgga actcgtcaag acaagtgtgg cagcagagga
                                                          200
151 gttttcaatt ttgatgaaca taagctggat actccgggca gacgccagca
                                                          250
152 teegettgtt gaaggeeace aagatetgeg tgagtggeaa aaacaacatg
                                                          300
153 aattcataca gctgtgtgag gtgcaactac acagaggcct tccaaagcca
                                                          350
154 gaccagacct tccggcggca aatggacatt ctcctatgta ggcttccctg
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                                                          450
155 tggagetgag cactetetat etcateageg cecataacat ceceaatget
156 aatatgaatg aggacagece ttetttgtet gtgaacttea cetegecagg
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                                                          550
157 ctqcctaaac cacqtaatga aatataaaaa gcagtgcact gaggcgggaa
                                                          600
158 qcctqtqqqa cccaqacatc actqcttgta aaaagaacga gaagatggtt
159 gaagtgaatt tcacaaccaa tccccttgga aacagataca cgattctcat
                                                          650
160 tcaacgggac acgacattgg ggttttctag agtgctggag aataaactga
                                                          700
                                                          750
161 tgaggacgtc tgtagccatc ccggtgactg aggagagtga aggtgcggtg
162 gttcagctga ccccatattt acatacctgc ggcaatgact gcatccgacg
                                                          800
163 cgaagggaca gttgtgcttt gctcagagac aagtgctccc atccctccag
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164 atgacaacag acgcatgctg ggaggctggc tgcctctctt cctggtgctg
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RAW SEQUENCE LISTING DATE: 10/30/2001 PATENT APPLICATION: US/09/778,971 TIME: 13:01:02

Input Set : A:\6138SEQ.txt

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166 aggaaggage acgaagacgt cettteetat ttecaceatg etectgeece 1000
167 tcattaaggt cctggtggtt tatccttctg agatatgttt ccatcacacc 1050
168 gtctgtcgct tcactgactt tcttcaaaac tactgcagaa gtgaggtcat 1100
169 ccttgaaaaa tggcagaaaa agaaaatcgc cgagatgggg ccggtacagt 1150
170 ggctgaccac tcagaagcaa gcggcagata aagtggtctt ccttcttccc 1200
171 agtgacgtcc cgaccctttg tgacagtgcc tgtggccaca atgagggcag 1250
172 cgccagggag aactctcagg atctgttccc tcttgccttt aacctctttt 1300
173 gtagtgattt cagcagccag acgcatctgc acaaatacct ggtggtctat 1350
174 cttgggggag cagacctcaa aggcgactat aatgccctga gtgtctgccc 1400
175 ccaatatcat ctcatgaagg acgccacagc tttccacaca gaacttctca 1450
176 aggctacgca gagcatgtca gtgaagaaac gctcacaagc ctgccatgat 1500
177 agetgtteae cettgtagte caeceggggg aatagagact etgaageett 1550
178 cctactctcc cttccaqtqa caaatqctqt qtqacqactc tgaaatgtqt 1600
179 gggagagget gtgtggaggt agtgetatgt acaaacttge tttaaaactg 1650
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181 ggatttatga agacaacaca gttacagaca ataatgagtg ggacctacat 1750
182 ttgggatata cccaaagctg ggtaatgatt atcactgaga accacgcact 1800
183 ctggccatga agtaatacgg cacttccctg tcaggctgtc tgtcaggttg 1850
184 gqtctqtctt qcactqccca tqctctatqc tqcacqtaga ccgttttgta 1900
185 acattttaat ctgttaatga ataatccgtt tgggaagctc tcaaaaaaaa 1950
186 aaaaaaaaa aaa
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189 <211> LENGTH: 2589
190 <212> TYPE: DNA
191 <213> ORGANISM: Unknown
193 <220> FEATURE:
195 <221> NAME/KEY: prim_transcript
196 <223> OTHER INFORMATION: cDNA of mouse Evi27
198 <400> SEOUENCE: 4
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200 gtgcaggagc gccctgcctc gagagccgac tattcagtgt ggctctgaga
                                                             100
201 cagggecate tecagagtgg atggtecaae acacacteae tecaggagae
                                                             150
202 ttgagggacc tccaagtgga actcgtcaag acaagtgtgg cagcagagga
                                                             200
203 gttttcaatt ttgatgaaca taagctggat actccgggca gacgccagca
                                                             250
204 tecgettgtt gaaggecace aagatetgeg tgagtggcaa aaacaacatg
                                                             300
205 aattcataca gctgtgtgag gtgcaactac acagaggcct tccaaagcca
                                                             350
206 gaccagacct tccggcggca aatggacatt ctcctatgta ggcttccctg
                                                             400
207 tggagctgag cactctctat ctcatcagcg cccataacat ccccaatgct
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208 aatatgaatg aggacageee ttetttgtet gtgaaettea eetegeeagg
209 gtgcactcgt gaaaacacag aagtaacgtc cggtgtattt ccagcagcta
                                                             550
210 aacaccaggc tctccggatt tcagctcctt tcccattaca atttcctcct
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                                                             650
211 qqqccaqaqq actcaqtcat tctgccaccc cagcctctgq cgtcgctttt
212 toatgacttt gtcaaactta cctagcttgt ttccattctg aaattgtctg
                                                             700
213 atgcttgctt cgtatgtaag ccggggatat gaggtttggg tatgaatccc
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                                                             800
214 acagagggca ctgaattctt ctcactatgg cctatctggg ctgtgtgaca
215 ttgttggtga gggtcgtgcc tactaggcat ctgggtatct accacctgga
                                                             850
216 gcttcatgtc tggaagaggc agaacctata tgtattgtca gctctcactt
217 ttgtttccgt gtcacctcct ggagactgtt tttgataaag gttgtactta
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/778,971

DATE: 10/30/2001 TIME: 13:01:02

Input Set : A:\6138SEQ.txt

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219 gttgtctctc atatctgaag taagtgtgtg tgtgtgtgtt ttgtgtgtgt 1050
220 gtgtgtgtgt gtgtgtgtac tgggcaaagg gttatacctt tactcaaatg 1100
221 taacaacttt cattcacatt cccaggctgc ctaaaccacg taatgaaata 1150
222 taaaaagcag tgcactgagg cgggaagcct gtgggaccca gacatcactg 1200
223 cttgtaaaaa gaacgagaag atggttgaag tgaatttcac aaccaatccc 1250
224 cttggaaaca gatacacgat tctcattcaa cgggacacga cattggggtt 1300
225 ttctagagtg ctggagaata aactgatgag gacgtctgta gccatcccgg 1350
226 tgactgagga gagtgaaggt gcggtggttc agctgacccc atatttacat 1400
227 acctgcggca atgactgcat ccgacgcgaa gggacagttg tgctttgctc 1450
228 agagacaagt gctcccatcc ctccagatga caacagacgc atgctgggag 1500
229 gctggctgcc tctcttcctg gtgctgctgg tggctgtgtg ggtgctggca 1550
230 gctgggatet acctaacttg gaggcaagga aggagcacga agacgtcett 1600
231 tectattice accategic tececitical taageticity giggittate 1650
232 cttctqaqat atqtttccat cacaccqtct gtcqcttcac tgactttctt 1700
233 caaaactact qcaqaaqtqa qqtcatcctt gaaaaatggc agaaaaagaa 1750
234 aatcgccgag atggggccgg tacagtggct gaccactcag aagcaagcgg 1800
235 cagataaagt ggtcttcctt cttcccagtg acgtcccgac cctttgtgac 1850
236 agtgcctgtg gccacaatga gggcagcgcc agggagaact ctcaggatct 1900
237 gttccctctt gcctttaacc tcttttgtag tgatttcagc agccagacgc 1950
238 atctgcacaa atacctggtg gtctatcttg ggggagcaga cctcaaaggc 2000
239 gactataatg ccctgagtgt ctgcccccaa tatcatctca tgaaggacgc 2050
240 cacagettte cacacagaae tteteaagge taegeagage atgteagtga 2100
241 agaaacgete acaageetge catgataget gttcaccett gtagtecace 2150
242 cgggggaata gagactctga agccttccta ctctcccttc cagtgacaaa 2200
243 tgctgtgtga cgactctgaa atgtgtggga gaggctgtgt ggaggtagtg 2250
244 ctatgtacaa acttgcttta aaactggagt ttgcaaagtc aacctgagca 2300
245 tacacgcctg aggctagtca ttggctggat ttatgaagac aacacagtta 2350
246 cagacaataa tgagtgggac ctacatttgg gatataccca aagctgggta 2400
247 atgattatca ctgagaacca cgcactctgg ccatgaagta atacggcact 2450
248 tecetgteag getgtetgte aggttgggte tgtettgeae tgeecatget 2500
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252 <210> SEQ ID NO: 5
253 <211> LENGTH: 502
254 <212> TYPE: PRT
255 <213> ORGANISM: Homo sapiens
257 <220> FEATURE:
259 <221> NAME/KEY: peptide
260 <223> OTHER INFORMATION: Human Evi27 protein
262 <400> SEQUENCE: 5
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265 Val Pro Arg Glu Pro Thr Val Gln Cys Gly Ser Glu Thr Gly Pro
266
                     20
267 Ser Pro Glu Trp Met Leu Gln His Asp Leu Ile Pro Gly Asp Leu
                                         40
                                                              45
269 Arg Asp Leu Arg Val Glu Pro Val Thr Thr Ser Val Ala Thr Gly
270
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VERIFICATION SUMMARY

DATE: 10/30/2001

PATENT APPLICATION: US/09/778,971

TIME: 13:01:03

Input Set : A:\6138SEQ.txt

Output Set: N:\CRF3\10302001\1778971.raw

L:103~M:258~W: Mandatory Feature missing, <222> not found for SEQ ID#:2

L:103 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2